

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> 891909,567A
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped	
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input checked="" type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <input type="checkbox"/> Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/909,567A

DATE: 12/12/2001
TIME: 11:40:36

Input Set : A:\DEX-214.txt
Output Set: N:\CRF3\12112001\I909567A.raw

Does Not Comply
Corrected Diskette Needed

Errors on pp. 2 & 4

3 <110> APPLICANT: Macina, Roberto A
4 Nair, Manoj
5 Chen, Seiyu
7 <120> TITLE OF INVENTION: Compositions and Methods Relating to Lung Specific
8 Genes
10 <130> FILE REFERENCE: DEX-0214
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/909,567A
C--> 13 <141> CURRENT FILING DATE: 2001-07-20
15 <150> PRIOR APPLICATION NUMBER: 60/219,834
16 <151> PRIOR FILING DATE: 2000-07-21
18 <160> NUMBER OF SEQ ID NOS: 56
20 <170> SOFTWARE: PatentIn Ver. 2.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 1449
24 <212> TYPE: DNA
25 <213> ORGANISM: Homo sapiens
27 <400> SEQUENCE: 1
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30 cagcctgaga gcccgggaca cggctgttta tttctgtgcg agagagccac cagcacccaa 180
31 ttactttgac tgctggagcc agggAACCT ggtcaccgtc tcctcagctt ccaccaagg 240
32 cccatcggtc ttccccctgg cggccctgctc caggagcacc tctggggca cagcggccct 300
33 gggctgcctg gtcaaggact acttccccga accggtgacg gtgtcgtgga actcaggcgc 360
34 cctgaccaggc ggcgtgcaca cttccggc tgcctacag tcctcaggac tctactccct 420
35 cagcagcgtg gtgacccgtc cttccagcag cttggcacc cagacctaca cctgcaacgt 480
36 gaatcacaag cccagcaaca ccaagggtgga caagagagtt gagctaaaaa ccccaacttgg 540
37 tgacacaact cacacatgcc cacgggtccc agagccaaa tcttggaca cacctcccc 600
38 atgcccacgg tgcccaagac ccaaattttg tgacacaccc ccccatgcc cacgggtccc 660
39 agcacctgaa ctccctggag gaccgtcagt cttccatccc ccccaaaaac ccaaggatac 720
40 ctttatgatt tcccgaccc ctgagggtcac gtgcgtggg gtggacgtga gccacgaaga 780
41 ccccgagggtc cagttcaagt ggtacgtgga cggcgtggag gtgcataatg ccaagacaaa 840
42 gccgcgggag gacgttca acagcacgtt ccgtgtggc acgtcctca cggccttgca 900
43 ccaggactgg ctgaacggca aggagtacaa gtcaagggtc tccaaacaaag ccctccca 960
44 ccccatcgag aaaaccatct caaaaaccaa aggacagccc cgagaaccac aggtgtacac 1020
45 cctgccccca tcccgggagg agatgaccaa gaaccagggtc acgttgcact gcctggtaa 1080
46 aggcttctac cccagcgaca tcgccgtgga gtggagagc acggggcagc cggagaacaa 1140
47 ctacaacacc acgcctccca tgctggactc cgacggctcc ttcttccctt acagcaagct 1200
48 caccgtggac aagagcaggt ggcacggg gaacatctt tcatgtccg tgatgcata 1260
49 ggctctgcac aaccgttca cgcagaagag cctctccctg tctccgggtt aatgagtgcg 1320
50 acggccggca agcccccgct ccccggtc tcggggtcg cgcaggatgc ttggcacgt 1380
51 ccccggtac atacttcccg ggcacccagc atggaaataa agcacccagc gctgccctgg 1440
52 gccccctgcg
55 <210> SEQ ID NO: 2
56 <211> LENGTH: 3825
57 <212> TYPE: DNA
58 <213> ORGANISM: Homo sapiens
60 <220> FEATURE:

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Input Set : A:\DEX-214.txt
Output Set: N:\CRF3\12112001\I909567A.raw

61 <221> NAME/KEY: unsure
62 <222> LOCATION: (428)
64 <220> FEATURE:
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66 <222> LOCATION: (430)
68 <220> FEATURE:
69 <221> NAME/KEY: unsure
70 <222> LOCATION: (1997)..(1998)
72 <400> SEQUENCE: 2
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74 cttaaaaaac acggccacccg tcaccaactt ttatcacaat agaatctact gcccgacgaa 120
75 cagaaaaccc tactaagaac gagcttttc agtccccctaa aaaggacagt tatgttgaac 180
76 ccccaccaag aaggcccatt tcgcaaaaat ctgaaattca cagagcaaacttccctt 240
77 ctccaccaggagtcgcctt gaacaacttgc tcaagactcaa agacaccact gcaaagttat 300
78 ccaaaggggc catcccatgt ccagcagcaa ccccggttcc aattgttagag aagaggtctg 360
79 aaatcatcat gtctcctgca acacttcgtc gtcaaattaa gatagaaact cgtggtaggg 420
W--> 80 actctccnan ctacaatcac aataccagta aatataaaatc atgctgctag tggttccttc 480
81 agagaatctg tggacgctca agaggaaatc agggaaagtgg agaagagagc tacttatgtt 540
82 cataaaagatg gactaaattc cactgatcac atggtgcccg acactgaaag ttatgtgca 600
83 gttgaatca tccgcaaggt tgcaatgcct cctcgccctgt cagagcacac acagagatat 660
84 gaagcggcca accgaactgt tcaaattggct gaaaatttcg tgaatgaccc taaaatgaa 720
85 ataaacagat ggttcaggaa atttggatcat gcccagttt ctgaagcaaa gtcaaataaga 780
86 agagtttatg caaagggaga aacaaaccat aacatacaac aagaaagtgc tacattttgt 840
87 aaggaggaat ttggatttaac atctttatgg aacacgagtt ttacagactt ttcttgcaaa 900
88 catcctagag aactgcgaga aaagattctt gttaaaggcgc ccaggatctg ctctgaaacc 960
89 aggtctctaa gtgaacattt ctcaggatcg gatgcattt agatcaaat ttttggatctg 1020
90 aagatgaaaaa cctcttcatac acatagctca gaagctggca aatctggctg tgacttcaag 1080
91 catgccccac caacctatga ggtatgtcatt gctggacata ttttagatat ctctgattca 1140
92 cctaaagaag taagaaaaaa tttcaaaatg acgtggcaag agatggaaatg agttttttaaa 1200
93 ggcctggat atgcaaccgc agatgttctt gcaactgaga tgagaaccac ctcccaagag 1260
94 gaatctgcatttataatgtca agtgcgtctt ccaagacaaatg gaaatatgtt tactttgtca 1320
95 aaagacagtt tatccaatgg agtgccttagt ggcagacaaatg cagaattttc ataagtccctg 1380
96 cttccgatgc caccattgca acagttaaact aatgttggaa aattatgtcat cacttcattgg 1440
97 acaaataatac tgtaaaccctc actttaaaca acttttcaaa tccaaaggaa attatgtga 1500
98 aggtttggaa cataaggcgc ataaagatag atggaaactgc aaaaacccaa gcatcgt 1560
99 ggactttatt cctaataatgg aacccaaatat gtgtaaaaat attgcagaaa acacccttg 1620
100 acctggagat cgtatgaac atttagatgc tgtaacagt gaaggcggaa ggaatgattt 1680
101 gagaaaaatggggaaaggaa gaaaattttaa agtcattttgg cctcccttcca aggagatccc 1740
102 taagaaaaacc ttacccttgc aggaagatgt caaaatgtt aaacctaaatg ggccacctga 1800
103 aatgacaacc ctgttatccc ctgaattttaa aatgttgcattt ctgttgcataatg atgtttagaaac 1860
104 tccagaaaaat aaaggacaaa gacaagatca ctttccattt ttgcacccctt atctacagtc 1920
105 caccatgtt tgcaatggaa aggtatgtt aggaatcaaa gaaatgaaaaa tgccctgaagg 1980
W--> 106 aagaaaaagat gaaaagnnng aaggaaggaa gaatgtgca gataaggccga gtgaagctga 2040
107 agacacaaag agtaacagga aaagtgcattt ggtatctttaat gacaacaata atgttgcattt 2100
108 gcagagtgc gaaaaggaga aaaatgaaaaa aactaaccatc actaatgttgc cagaagtttt 2160
109 acaggttact aacactgttgc atggatgttgc gccagaaaaat cttaaagaaa atttgcataatg 2220
110 gaataataat aacaattatgt tagcaatgttgc atatcttgcattt aatttgcaggc agaagacatc 2280
111 tatttttagaa ttcttgcattt tattaccctt ttgcagttgc gcaaatgaca ctgcaatgtca 2340
112 atatgaaattt gagaagttag aaaatatac tagaatctca gagttacttgc gtatatttgc 2400

Please explain what residue represents in L2237. See error summary sheet

file:///C:/CrF3/Outhold/VsrI909567A.htm

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/909,567A

DATE: 12/12/2001
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Input Set : A:\DEX-214.txt
Output Set: N:\CRF3\12112001\I909567A.raw

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Input Set : A:\DEX-214.txt
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165 actttgagca gttagaagtg ctcttttagaa gttgtctgtg ggtgatgtta ctgttagtggt 1260
 166 ctcagggaaa ggattgtcca gttactttag ggggttttg gtgggggttt tccccctgtg 1320
 167 aaaacttaact ttgcccctag tctggctgtc gctaggactt ctgaggagca atgggacatg 1380
 168 agtgtccctg tatctgcgcc actgccgcaa gggaaaccc aggaaccaggc acctggaggc 1440
 169 caggatagcc aagccctggg tgagcgagag gctggagaac acaggagctc acccaggc 1500
 170 gctgcccaac catggccac tctgaaacaga cttcagtcct ctgttttg ttcataagcc 1560
 171 gttgagacat ctgtatggact tggcttaggc cctgtggga catcccacgt gtgatccctt 1620
 172 tcactccatc aggacaccag gactgtcctt agggaaaatgt ctttgagatg gcagcaggag 1680
 173 tcataatttc tggatgtgtg tttcgaaag ccgtgtgtc ctgcctcagc acaaagaccc 1740
 174 agtgtcattt gctccctgtt tcctgtgcc actccagaac ctcagcagat ctgagccacc 1800
 175 gcctgccagt gtgagaggcg gccactttca tggcagctca tcaggcgcag ggcggcagac 1860
 176 agcttcccaag caggccctag agcccgccct gggccatga tggagggcgg ccgcagcc 1920
 177 agggcctgcc catccagaag gactcccca gggctgggg gaggagaccc ttggaaaagt 1980
 178 cctctcttcc cagctcctga ttctggatct gagattctca gatcacaggc ccctgtgtc 2040
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 181 tcaggcctgg gccatgtgg tggattttctt aaagctggag aaaggaagaa ttgtgcctt 2220
 182 catattactt gagcttaaac tgacaacctg gatgtaaata ggaccccttc tactggttt 2280
 183 ttaataaaag ttctatgtga tttttaaaaa aaaaa 2315
 186 <210> SEQ ID NO: 4
 187 <211> LENGTH: 300
 188 <212> TYPE: DNA
 189 <213> ORGANISM: Homo sapiens
 190 <220> FEATURE: *Same error*
 191 <221> NAME/KEY: unsure
 192 <222> LOCATION: (8)
 193 <400> SEQUENCE: 4
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 198 agtgcctccca agaggagccca gggctggctg cagtggaggcc ccaggaggtt ctccagggc 180
 199 gtcctgcttc agctcaaggg gcttaggaata ggggaaacga tgcaggaaag ccaatggccc 240
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 204 <211> LENGTH: 4347
 205 <212> TYPE: DNA
 206 <213> ORGANISM: Homo sapiens
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 211 taagcacaag aaatgaaaat aagctgttcc ctaaacatcc tcattttatg cggcaaaagc 180
 212 ggcctgtat caccggcccc gttggctttc gggaggggaga ggatctgtcc aagaagaatc 240
 213 caattgccaa gatacatttct gatcttgcag aagaaagagg actcaaaat acttacaaat 300
 214 acactggaaa agggattaca gagccaccc ttgttatatt tgcattttaaac aaagataactg 360
 215 gagaactgaa tggatgttcc attcttgcac gagaagaaac accatttttt ctgctaaacag 420
 216 gttacgtttt ggtatgttcc gggaaacatg tagagaaacc ctttagagctc cgcattaaagg 480
 217 ttcttgcata caatgacaac gaaccagggt tcacacaggta tgcattttttt gggctgttg 540
 218 aagagttgag tgcagcacat actcttgcata tgaaaatcaa tgcaacagat gcaatgtgac 600
 219 ccaataccctt gaattcgaaa atttcctata gaatcgatcc tctggaggcct gcttatccctc 660
 220 cagtgttata cttaaataaa gatacaggag agatttatac aaccagggtt accttggaca 720

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Input Set : A:\DEX-214.txt
Output Set: N:\CRF3\12112001\I909567A.raw

221 gagaggaaca cagcagctac actttgacag tagaaagcaag agatggcaat ggagaagtt 780
 222 cagacaaacc tggaaaacaa gctcaagttc agattcgat tttggatgtc aatgacaata 840
 223 tacctgttagt agaaaataaa gtgcttgaag ggatgggtga agaaaatcaa gtcaacgtag 900
 224 aagttacgctg cataaaagtg ttgcgtgcag atgaaatagg ttctgataat tggctggcaa 960
 225 attttacatt tgcattcagg aatgaaggag gttatttcca catagaaaca gatgctcaa 1020
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 227 acttcagtgt tattgtcgat aataaaggcag cttttcacaa gtcgatttgg agtaaataca 1140
 228 agcctacacc cattccatc aaggtaaaag tgaaaaatgt gaaagaaggc attcatttt 1200
 229 aaagcagcgt catctcaatt tatgttagcg agagcatgga tagatcaagc aaaggccaa 1260
 230 taatttggaaa tttcaagct ttgtatgagg acactggact accagccat gcaagatatg 1320
 231 taaaattaga agatagagat aattggatct ctgtggattc tgcacatct gaaattaaac 1380
 232 ttgcaaaact tcctgattt gaatcttagat atgttcaaaa tggcacatc actgtaaaga 1440
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 237 aagaaagtac cagtgtgctg ctgcacacaa gtgagaaaaa gcttgggaga agtggaaattc 1740
 238 agttcctgtat ttcagacaaat cagggtttt gttgtcctgaa aagcagggtc cttacactca 1800
 239 cagttgtga gtgtcgat ggcagcggct gcaggaaagc acagcatgac tcctatgtgg 1860
 240 gcctgggacc cgcagcaatt ggcgtcatgat tttggccctt tctgtctgtg ctattggatc 1920
 241 cactttact gctgtatgtc cattgcggaa agggcgccaa aggctttacc cccataccgt 1980
 242 gcaccataga gatgtcgat ctttggataat atgaaggagc accacctgaa gacaagggtgg 2040
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 246 ctacccaggat tacagggggcc acaggcgctt tcatgaccatc tgaaaccacg aagaccgca 2280
 247 gggccacagg ggctccaga gacatggccg gagtcaggc agtcgtgtt gcactgaacg 2340
 248 aagaattttt aagaattat ttcactgata aagccgcctc ttacactgag gaagatgaaa 2400
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 260 aagatgttac ttatgttgcgaa ggggggttttccatc ttttttttttccatc ttttttttttccatc 3120
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 265 agcatagacat ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc 3420
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 267 agacacacacg agacacacatc acattgtatgttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc 3540
 268 ggaccacact gtttgcgttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc 3600
 269 atagatgttac ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc ttttttttttccatc 3660

VERIFICATION SUMMARY
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Input Set : A:\DEX-214.txt
Output Set: N:\CRF3\12112001\I909567A.raw

L:12 M:270 C: Current Application Number differs, Replaced Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:80 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:2
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:106 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:2
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:196 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:4
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:338 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:445 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:445 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:446 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:447 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:447 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:448 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:452 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:453 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:954 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:17
L:954 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:982 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:18
L:982 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1175 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1176 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1177 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1178 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1179 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1180 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:19
L:1180 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1399 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1400 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1401 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1402 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1403 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/909,567A

DATE: 12/12/2001

TIME: 11:40:37

Input Set : A:\DEX-214.txt

Output Set: N:\CRF3\12112001\I909567A.raw

L:1404 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22
L:1404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1738 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:38
L:1738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38